

***In the claims:***

Please amend claims 23, 25, 27, 29, 33-36, 39, 40, 43, 44, 47, 48, 60-67, 72-74, 77, 78, 81, 82, 85 and 86 as follows.

*D<sup>2</sup>* 3. ~~25.~~ (Once Amended) The polypeptide of claim ~~22~~<sup>3</sup>, wherein the amino acid sequence is fused to a heterologous polypeptide.

*D<sup>3</sup>* 5. ~~25.~~ (Once Amended) The polypeptide of claim ~~24~~<sup>4</sup>, wherein the amino acid sequence is fused to a heterologous polypeptide.

*D<sup>4</sup>* 7. ~~27.~~ (Once Amended) The polypeptide of claim ~~26~~<sup>6</sup>, wherein the amino acid sequence is fused to a heterologous polypeptide.

*D<sup>5</sup>* 9. ~~28.~~ (Once Amended) The polypeptide of claim ~~28~~<sup>8</sup>, wherein the amino acid sequence is fused to a heterologous polypeptide.

*D<sup>6</sup>* 10. ~~35.~~ (Once Amended) The polypeptide of claim ~~22~~<sup>21</sup>, wherein the amino acid sequence is fused to a heterologous polypeptide.

*D<sup>7</sup>* 11. ~~34.~~ (Twice Amended) An isolated polypeptide comprising a first amino acid sequence that is at least 95% identical to a second amino acid sequence selected from the group consisting of:

(a) amino acids 1 to 381 of SEQ ID NO:2;

D<sup>1</sup>  
cont'd

- (b) amino acids 2 to 381 of SEQ ID NO:2;
- (c) amino acids 25 to 381 of SEQ ID NO:2; and
- (d) a polypeptide fragment of SEQ ID NO:2,

wherein said polypeptide or polypeptide fragment stimulates cellular proliferation.

D<sup>8</sup>

12. <sup>35.</sup> (Once Amended) The polypeptide of claim <sup>11</sup>~~34~~, wherein said second amino acid sequence is (a).

D<sup>9</sup>

13. <sup>36.</sup> (Once Amended) The polypeptide of claim <sup>12,</sup>~~35~~, wherein the amino acid sequence is fused to a heterologous polypeptide.

D<sup>9</sup>

14. <sup>39.</sup> (Once Amended) The polypeptide of claim <sup>11</sup>~~34~~, wherein said second amino acid sequence is (b).

D<sup>9</sup>

15. <sup>40.</sup> (Once Amended) The polypeptide of claim <sup>14,</sup>~~39~~, wherein the amino acid sequence is fused to a heterologous polypeptide.

D<sup>10</sup>

16. <sup>42.</sup> (Once Amended) The polypeptide of claim <sup>11</sup>~~34~~, wherein said second amino acid sequence is (c).

D<sup>10</sup>

17. <sup>44.</sup> (Once Amended) The polypeptide of claim <sup>16,</sup>~~42~~, wherein the amino acid sequence is fused to a heterologous polypeptide.

*D* 18. (Once Amended) The polypeptide of claim ~~34~~<sup>11</sup>, wherein said second amino acid sequence is (d).

*D* 19. ~~48.~~ (Once Amended) The polypeptide of claim ~~47~~<sup>18</sup>, wherein the amino acid sequence is fused to a heterologous polypeptide.

*D* 20. ~~60.~~ (Once Amended) The polypeptide of claim ~~59~~<sup>20</sup>, wherein said amino acid sequence is (a).

*D* 21. ~~61.~~ (Once Amended) The polypeptide of claim ~~60~~<sup>21</sup>, wherein the amino acid sequence is fused to a heterologous polypeptide.

*D* 22. ~~62.~~ (Once Amended) The polypeptide of claim ~~59~~<sup>20</sup>, wherein said amino acid sequence is (b).

*D* 23. ~~63.~~ (Once Amended) The polypeptide of claim ~~62~~<sup>23</sup>, wherein the amino acid sequence is fused to a heterologous polypeptide.

*D* 24. ~~64.~~ (Once Amended) The polypeptide of claim ~~59~~<sup>20</sup>, wherein said amino acid sequence is (c).

*D* 25. ~~65.~~ (Once Amended) The polypeptide of claim ~~64~~<sup>25</sup>, wherein the amino acid sequence is fused to a heterologous polypeptide.

*D<sup>12</sup> cont'd* 27.

66. (Once Amended) The polypeptide of claim 59, wherein said amino acid sequence is (d).

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67. (Once Amended) The polypeptide of claim 68, wherein the amino acid sequence is fused to a heterologous polypeptide.

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72. (Twice Amended) An isolated polypeptide comprising a first amino acid sequence that is at least 95% identical to a second amino acid sequence selected from the group consisting of:

(a) the amino acid sequence of the full-length polypeptide encoded by the human cDNA contained in ATCC Deposit Number 75904;

(b) the amino acid sequence of the full-length polypeptide, lacking the N-terminal methionine, encoded by the human cDNA contained in ATCC Deposit Number 75904;

(c) the amino acid sequence of the mature polypeptide encoded by the human cDNA contained in ATCC Deposit Number 75904; and

(d) a polypeptide fragment of the polypeptide encoded by the human cDNA contained in ATCC Deposit Number 75904;

wherein said polypeptide or polypeptide fragment stimulates cellular proliferation.

*D<sup>14</sup>* 30

73. (Once Amended) The polypeptide of claim 29, wherein said second amino acid sequence is (a).

D<sup>14</sup> 31.  
74. (Once Amended) The polypeptide of claim ~~73~~<sup>30</sup>, wherein the amino acid sequence is fused to a heterologous polypeptide.

D<sup>15</sup> 32.  
77. (Once Amended) The polypeptide of claim ~~72~~<sup>29</sup>, wherein said second amino acid sequence is (b).

D<sup>16</sup> 33.  
78. (Once Amended) The polypeptide of claim ~~77~~<sup>32</sup>, wherein the amino acid sequence is fused to a heterologous polypeptide.

34.  
81. (Once Amended) The polypeptide of claim ~~72~~<sup>29</sup>, wherein said second amino acid sequence is (c).

D<sup>17</sup> 35.  
82. (Once Amended) The polypeptide of claim ~~81~~<sup>34</sup>, wherein the amino acid sequence is fused to a heterologous polypeptide.

D<sup>17</sup> 36.  
83. (Once Amended) The polypeptide of claim ~~72~~<sup>29</sup>, wherein said second amino acid sequence is (d).

D<sup>17</sup> 37.  
84. (Once Amended) The polypeptide of claim ~~83~~<sup>36</sup>, wherein the amino acid sequence is fused to a heterologous polypeptide.